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## ALIGNMENTS

## RESULT 1

US-09-325-256-24

; Sequence 24, Application US/09325256

; Patent No. 6444793

; GENERAL INFORMATION:

; APPLICANT: PEPINSKY, R. BLAKE

; APPLICANT: BAKER, DARREN P.

; APPLICANT: WEN, DINGYI

; APPLICANT: WILLIAMS, KEVIN P.

; APPLICANT: GARGER, ELLEN A.

; APPLICANT: TAYLOR, FREDERICK R.

; APPLICANT: GALDES, ALPHONSE

; APPLICANT: PORTER, JEFFREY

; TITLE OF INVENTION: HYDROPHOBICALLY-MODIFIED PROTEIN COMPOSITIONS AND

; TITLE OF INVENTION: METHODS

; FILE REFERENCE: BIV-067.01

; CURRENT APPLICATION NUMBER: US/09/325,256

; CURRENT FILING DATE: 1999-06-03

; PRIOR APPLICATION NUMBER: 60/099,800

; PRIOR FILING DATE: 1998-09-10

; PRIOR APPLICATION NUMBER: 60/078,935

; PRIOR FILING DATE: 1998-03-20

; PRIOR APPLICATION NUMBER: 60/089,685

; PRIOR FILING DATE: 1998-06-17

; PRIOR APPLICATION NUMBER: 60/067,423

; PRIOR FILING DATE: 1997-12-03

; PRIOR APPLICATION NUMBER: PCT/US98/25676

; PRIOR FILING DATE: 1998-12-03

; NUMBER OF SEQ ID NOS: 31

; SOFTWARE: PatentIn Ver. 2.1

; SEQ ID NO 24

; LENGTH: 396

; TYPE: PRT

; ORGANISM: Homo sapiens

US-09-325-256-24

GenCore version 5.1.8  
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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Perfect score: 2088  
Sequence: 1 MALLTNLLPLCCLALLALPA.....PTGMHWYSRLLYRLAEELLG 396

Scoring table: BLOSUM62  
Gapop 10.0 , Gapext 0.5

Searched: 56051 seqs, 11137335 residues

Total number of hits satisfying chosen parameters: 56051

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

#### SUMMARIES

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5	84	4.0	227	6	US-10-953-349-34128
6	80.5	3.9	448	7	US-11-293-697-3667
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